



IFW16

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/920,137B

DATE: 10/01/2004
TIME: 10:46:27

Input Set : A:\Cen 0250 SEQ LIST 07-09-01.txt
Output Set: N:\CRF4\10012004\I920137B.raw

5 <110> APPLICANT: Giles-Komar, Jill
6 David Shealy
7 David Knight
8 Bernie Scallon
9 George Heavner
11 <120> TITLE OF INVENTION: ANTI-TNF ANTIBODIES, COMPOSITIONS, METHODS AND USES
13 <130> FILE REFERENCE: CEN0250
C--> 15 <140> CURRENT APPLICATION NUMBER: US/09/920,137B
C--> 15 <141> CURRENT FILING DATE: 2001-08-01
15 <160> NUMBER OF SEQ ID NOS: 15
17 <170> SOFTWARE: PatentIn Ver 3.1

ERRORED SEQUENCES

19 <210> SEQ ID NO: 1
20 <211> LENGTH: 5
21 <212> TYPE: PRT
22 <213> ORGANISM: Homo sapiens
W--> 23 <400> SEQUENCE: 1
25 Arg Tyr Thr Met His
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28 <210> SEQ ID NO: 2
29 <211> LENGTH: 17
30 <212> TYPE: PRT
31 <213> ORGANISM: Homo sapiens
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34 Val Ile Ser Phe Asp Gly Ser Asn Lys Tyr Tyr Val Asp Ser Val Lys
E--> 35 1 1 5 ✓ 10 10 15 ✓
37 <210> SEQ ID NO: 3
38 <211> LENGTH: 10
39 <212> TYPE: PRT
40 <213> ORGANISM: Homo sapiens
W--> 41 <400> SEQUENCE: 3
43 Glu Ala Arg Gly Ser Tyr Ala Phe Asp Ile
E--> 44 1 X 5 ✓ 10 ✓
64 <210> SEQ ID NO: 6
65 <211> LENGTH: 10
66 <212> TYPE: PRT
67 <213> ORGANISM: Homo sapiens
W--> 68 <400> SEQUENCE: 6
70 Gln Glp Arg Ser Asn Trp Pro Pro Phe Thr
E--> 71 1 X 5 5 10 ✓

U.S. Non-Comply
Corrected Diskette Needed

(PG. 1-3)

Pls see item
3 on error
summary
sheet.

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Input Set : A:\Cen 0250 SEQ LIST 07-09-01.txt
Output Set: N:\CRF4\10012004\I920137B.raw

192 <210> SEQ ID NO. 13
193 <211> LENGTH: 32 *Found 42*
194 <212> TYPE: DNA
195 <213> ORGANISM: Homo sapiens
W--> 196 <400> SEQUENCE: 13
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32 42

VERIFICATION SUMMARY

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Input Set : A:\Cen 0250 SEQ LIST 07-09-01.txt
Output Set: N:\CRF4\10012004\I920137B.raw

L:15 M:270 C: Current Application Number differs, Replaced Current Application No
L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:23 M:283 W: Missing Blank Line separator, <400> field identifier
L:26 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1 ✓
L:32 M:283 W: Missing Blank Line separator, <400> field identifier
L:35 M:252 E: No. of Seq. differs, <211> LENGTH:Input:17 Found:16 SEQ:2 ✓
L:41 M:283 W: Missing Blank Line separator, <400> field identifier ✓
L:44 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3 ✓
L:50 M:283 W: Missing Blank Line separator, <400> field identifier
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L:68 M:283 W: Missing Blank Line separator, <400> field identifier
L:71 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6 ✓
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L:108 M:283 W: Missing Blank Line separator, <400> field identifier
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L:188 M:283 W: Missing Blank Line separator, <400> field identifier
L:196 M:283 W: Missing Blank Line separator, <400> field identifier
L:198 M:254 E: No. of Bases conflict, LENGTH:Input:32 Counted:42 SEQ:13 ✓
L:198 M:252 E: No. of Seq. differs, <211> LENGTH:Input:32 Found:42 SEQ:13 ✓
L:204 M:283 W: Missing Blank Line separator, <400> field identifier
L:212 M:283 W: Missing Blank Line separator, <400> field identifier